

RAW SEQUENCE LISTING
ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/627,165

Source: OIPE

Date Processed by STIC: 4/16/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/627,165

DATE: 04/16/2001
TIME: 11:22:51

Input Set : A:\CRFascidoc.txt.txt
Output Set: N:\CRF3\04162001\I627165.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: KIM, Jong-Bae
5 <120> TITLE OF INVENTION: CRUDE EXTRACT FROM Viscum album coloratum, AND PROTEINS
6 AND LECTINS ISOLATED THEREFROM
8 <130> FILE REFERENCE: Korean Mistletoe Lectin
10 <140> CURRENT APPLICATION NUMBER: 09/627,165
11 <141> CURRENT FILING DATE: 2000-07-27
13 <160> NUMBER OF SEQ ID NOS: 16
15 <210> SEQ ID NO: 1

Delete circled item below.
because no n's exist in this
sequence

This is a nucleic acid sequence

16 <211> LENGTH: 762
17 <212> TYPE: DNA
18 <213> ORGANISM: Viscum album coloratum
19 <221> NAME/KEY: misc_feature
20 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
21 <400> SEQUENCE: 1

↓ FYI!
"n" can only
represent
a single
nucleotide.
n ≠ an amino
acid

22 tacgagagggc taagactcag agttacgcat caaaccacgg gcgacgaata ttcccggttc 60
23 atcacgcttc tccgagatta tgtctcaagc ggaagctttt ccaatgagat accactcttg 120
24 cgtcagtcta cgatccccgt ctccgatgag caaagatttg tggttggtgga actcaccaat 180
25 cagggggggag actcgtacac ggccgccatc gacgttacta acctgtacgt ggtggcttac 240
26 caagcaggcg accaatccta ctttttgccg gacgcaccag acggcgcgga aaggcatctc 300
27 ttcaccggca ccaccagatc ctccctccca ttcaccggaa gctacacaga tctggagcga 360
28 ttcgccggtc atagggacca gatccctctg ggtagagagg aactcattca atccgtctcg 420
29 gcccttcgtt ttccggggcag caacactcgt gcccaagctc gttcctttat catcctcatt 480
30 cagatgatct ccgaggccgc cagattcaat cccatcttat ggagggtcgc ccaatacatt 540
31 agcagtggggg ggtcattttct gccagacacg tacattctcc agctggagac gagttggggg 600
32 caacaatcca cgcaagttca gcactcgacg gatggcggtt ttaataaccc aattcggttg 660
33 actatatcca ctggtgtctt cgtgacgttg agcaatgttc gcgacgtgat cgccagctta 720
34 gcgatcatgt tgtttgtatg cgaggaccgg ccatcttctc ct 762

FYI "n" does not belong in an
amino
acid
sequence

37 <210> SEQ ID NO: 2
38 <211> LENGTH: 254
39 <212> TYPE: PRT
40 <213> ORGANISM: Viscum album coloratum
41 <221> NAME/KEY: misc_feature
42 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
44 <400> SEQUENCE: 2

use Xaa
in an amino
acid sequence

45 Tyr Glu Arg Leu Arg Leu Arg Val Thr His Gln Thr Thr Gly Asp Glu
46 1 5 10 15
48 Tyr Phe Arg Phe Ile Thr Leu Leu Arg Asp Tyr Val Ser Ser Gly Ser
49 20 25 30
51 Phe Ser Asn Glu Ile Pro Leu Leu Arg Gln Ser Thr Ile Pro Val Ser
52 35 40 45
54 Asp Ala Gln Arg Phe Val Leu Val Glu Leu Thr Asn Gln Gly Gly Asp
55 50 55 60
57 Ser Ile Thr Ala Ala Ile Asp Val Thr Asn Leu Tyr Val Val Ala Tyr
58 65 70 75 80
60 Gln Ala Gly Asp Gln Ser Tyr Phe Leu Arg Asp Ala Pro Asp Gly Ala
61 85 90 95
63 Glu Arg His Leu Phe Thr Gly Thr Thr Arg Ser Ser Leu Pro Phe Thr

FYI: no Xaals
exist in
this sequence

Delete circled item

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/627,165

DATE: 04/16/2001
TIME: 11:22:51

Input Set : A:\CRFascidoc.txt.txt
Output Set: N:\CRF3\04162001\I627165.raw

```

64          100          105          110
66 Gly Ser Tyr Thr Asp Leu Glu Arg Phe Ala Gly His Arg Asp Gln Ile
67          115          120          125
69 Pro Leu Gly Arg Glu Glu Leu Ile Gln Ser Val Ser Ala Leu Arg Phe
70          130          135          140
72 Pro Gly Ser Asn Thr Arg Ala Gln Ala Arg Ser Phe Ile Ile Leu Ile
73 145          150          155          160
75 Gln Met Ile Ser Glu Ala Ala Arg Phe Asn Pro Ile Leu Trp Arg Ala
76          165          170          175
78 Arg Gln Tyr Ile Ser Ser Gly Gly Ser Phe Leu Pro Asp Thr Tyr Ile
79          180          185          190
81 Leu Gln Leu Glu Thr Ser Trp Gly Gln Gln Ser Thr Gln Val Gln His
82          195          200          205
84 Ser Thr Asp Gly Val Phe Asn Asn Pro Ile Arg Leu Thr Ile Ser Thr
85          210          215          220
87 Gly Val Phe Val Thr Leu Ser Asn Val Arg Asp Val Ile Ala Ser Leu
88 225          230          235          240
90 Ala Ile Met Leu Phe Val Cys Glu Asp Arg Pro Ser Ser Ser
91          245          250

94 <210> SEQ ID NO: 3
95 <211> LENGTH: 762
96 <212> TYPE: DNA
97 <213> ORGANISM: Viscum album coloratum
98 <221> NAME/KEY: misc_feature
99 <223> OTHER INFORMATION: (any set containing "n"=X, X=any amino acid)
101 <400> SEQUENCE: 3
102 tacgagagggc taagactcag agttacgcat caaaccacgg gcgaccaata tttcaagttc 60
103 atcacgcttc tccgagatca tgtctcaagc ggaagcttgt ccaatcaaata accactcttg 120
104 cggcagtgcta ctgtccccgt ctoggatacg cagagatttg tggttggtgga actcagcaat 180
105 cagggggggag actcgatcac ggcggccatc gacgttacca atctgtacgt ggtggcttac 240
106 caagcaggca accaatccta ctttttgccg gacgcacctc gcggcgcgga aacgtatctc 300
107 ttcaccggca ccaccgatc ctctctccca ttcaacggaa gctaccctga tctggagcga 360
108 tacgcccggac atagggacca gatccctctc ggtatagacc aactcattca atccgtctcg 420
109 gcccttcggt ttccgggcag caacactcgt gcccaagctc gttcctttat catcctcatt 480
110 cagatgatct cggaggccgc cagattcaat cccatcttat ggagggtctg ccaatacatt 540
111 agcagtgggg ggtcatttct gccagacacg tacattctcc agctggagac gagttggggg 600
112 caacaatcca cgcaagttca gcactcgacg gatggcggtt ttaataaacc aattcggttg 660
113 actatatcca ctggtgtctt cgtgacgttg agcaatgttc gcgacgtgat cgccagcyta 720
114 gcgatcatgt tgtttgtatg cgaggaccgg ccatcttctt ct 762
117 <210> SEQ ID NO: 4
118 <211> LENGTH: 254
119 <212> TYPE: PRT
120 <213> ORGANISM: Viscum album coloratum
121 <221> NAME/KEY: misc_feature
122 <223> OTHER INFORMATION: (any set containing "n"=X, X=any amino acid)
124 <400> SEQUENCE: 4
125 Tyr Glu Arg Leu Arg Leu Arg Val Thr His Gln Thr Thr Gly Asp Gln
126 1          5          10          15
128 Tyr Phe Lys Phe Ile Thr Leu Leu Arg Asp His Val Ser Ser Gly Ser

```

Delete circled item:
This is a nucleic acid sequence.

"n" can only represent a single nucleotide

(No n's exist in this sequence)

Insert 22207 wherever 22217, 22227, or 22237 is shown

Xaa Xaa = any amino acid

delete "n"=X

This is an amino acid sequence -

do not show any statement pertaining to n's

RAW SEQUENCE LISTING
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Input Set : A:\CRFascidoc.txt.txt
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```

129          20          25          30
131 Leu Ser Asn Gln Ile Pro Leu Leu Arg Gln Ser Thr Val Pro Val Ser
132          35          40          45
134 Asp Thr Gln Arg Phe Val Leu Val Glu Leu Ser Asn Gln Gly Gly Asp
135          50          55          60
137 Ser Ile Thr Ala Ala Ile Asp Val Thr Asn Leu Tyr Val Val Ala Tyr
138 65          70          75          80
140 Gln Ala Gly Asn Gln Ser Tyr Phe Leu Arg Arg Ala Pro Arg Gly Ala
141          85          90          95
143 Glu Thr Tyr Leu Phe Thr Gly Thr Thr Arg Ser Ser Leu Pro Phe Asn
144          100          105          110
146 Gly Ser Tyr Pro Asp Leu Glu Arg Tyr Ala Gly His Arg Asp Gln Ile
147          115          120          125
149 Pro Leu Gly Ile Asp Gln Leu Ile Gln Ser Val Ser Ala Leu Arg Phe
150 130          135          140
152 Pro Gly Ser Asn Thr Arg Ala Gln Ala Arg Ser Phe Ile Ile Leu Ile
153 145          150          155          160
155 Gln Met Ile Ser Glu Ala Ala Arg Phe Asn Pro Ile Leu Trp Arg Ala
156          165          170          175
158 Arg Gln Tyr Ile Ser Ser Gly Gly Ser Phe Leu Pro Asp Thr Tyr Ile
159          180          185          190
161 Leu Gln Leu Glu Thr Ser Trp Gly Gln Gln Ser Thr Gln Val Gln His
162          195          200          205
164 Ser Thr Asp Gly Val Phe Asn Asn Pro Ile Arg Leu Thr Ile Ser Thr
165 210          215          220
167 Gly Val Phe Val Thr Leu Ser Asn Val Arg Asp Val Ile Ala Ser Xaa
168 225          230          235          240
170 Ala Ile Met Leu Phe Val Cys Glu Asp Arg Pro Ser Ser Ser
171          245          250
174 <210> SEQ ID NO: 5
175 <211> LENGTH: 768
176 <212> TYPE: DNA
177 <213> ORGANISM: Viscum album coloratum
178 <221> NAME/KEY: misc_feature
179 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
181 <400> SEQUENCE: 5
182 tacgagaggc taagactcag agttacgcat caaaccacgg gcgacgaata tttccggttc 60
183 atcaagcttc tccgagactc tgtctcaagc ggaagctttt ccaatgacat accgctcctg 120
184 cctccgtcaa tcccgggtct ctctgcgcag agatttgtgt tgggtggaact cacaatcag 180
185 ttgggaaagt gggaagactc gatcacggcc gccatcgacg ttaccaatct gtacgtggtg 240
186 gcttaccaag caggcgacca atcctacttt ttgcgcgacg caccagacgg cgcggaaagg 300
187 catctcttca ccggcaccac cagatcctct ctctctttca acggaagcta cgctgatctg 360
188 gagcggtacg ccggacatag ggaccggatc cctctgggta gagagccact catacgatcc 420
189 gtctcggcgc ttgattatcc cggcggcagc acgcgcgccc aagccagttc cattattatc 480
190 gtcattcaga tgatctccga ggcggccaga ttcaatocca tcctatggag ggctcgccaa 540
191 tacattaaca gtggggtgtc atatcttcca gacgtgtaca tgctggagct ggaggcgagt 600
192 tggggccaac aatcgaccca agtccagcag tcgaccgatg gcgtttttta taaccaatt 660
193 cggttggtga tatccaccgg caacttcgtg tggttgagca atgttcgcga cgtgatcgcc 720
194 agcttgggga tcatggtgtt tgtatgcagg gaccggtcat cttcccct 768

```

no n's exist in this sequence

delete circled item

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/627,165

DATE: 04/16/2001
TIME: 11:22:51

Input Set : A:\CRFascidoc.txt.txt
Output Set: N:\CRF3\04162001\I627165.raw

```

197 <210> SEQ ID NO: 6
198 <211> LENGTH: 256
199 <212> TYPE: PRT
200 <213> ORGANISM: Viscum album coloratum
201 <221> NAME/KEY: misc_feature
202 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
204 <400> SEQUENCE: 6
205 Tyr Glu Arg Leu Arg Leu Arg Val Thr His Gln Thr Thr Gly Asp Glu
206 1 5 10 15
208 Tyr Phe Arg Phe Ile Lys Leu Leu Arg Asp Ser Val Ser Ser Gly Ser
209 20 25 30
211 Phe Ser Asn Asp Ile Pro Leu Leu Pro Pro Ser Ile Pro Val Ser Ser
212 35 40 45
214 Ala Gln Arg Phe Val Leu Val Glu Leu Thr Asn Gln Leu Gly Lys Trp
215 50 55 60
217 Glu Asp Ser Ile Thr Ala Ala Ile Asp Val Thr Asn Leu Tyr Val Val
218 65 70 75 80
220 Ala Tyr Gln Ala Gly Asp Gln Ser Tyr Phe Leu Arg Asp Ala Pro Asp
221 85 90 95
223 Gly Ala Glu Arg His Leu Phe Thr Gly Thr Thr Arg Ser Ser Leu Pro
224 100 105 110
226 Phe Asn Gly Ser Tyr Ala Asp Leu Glu Arg Tyr Ala Gly His Arg Asp
227 115 120 125
229 Arg Ile Pro Leu Gly Arg Glu Pro Leu Ile Arg Ser Val Ser Ala Leu
230 130 135 140
232 Asp Tyr Pro Gly Gly Ser Thr Arg Ala Gln Ala Ser Ser Ile Ile Ile
233 145 150 155 160
235 Val Ile Gln Met Ile Ser Glu Ala Ala Arg Phe Asn Pro Ile Leu Trp
236 165 170 175
238 Arg Ala Arg Gln Tyr Ile Asn Ser Gly Val Ser Tyr Leu Pro Asp Val
239 180 185 190
241 Tyr Met Leu Glu Leu Glu Ala Ser Trp Gly Gln Gln Ser Thr Gln Val
242 195 200 205
244 Gln Gln Ser Thr Asp Gly Val Phe Asn Asn Pro Ile Arg Leu Gly Ile
245 210 215 220
247 Ser Thr Gly Asn Phe Val Trp Leu Ser Asn Val Arg Asp Val Ile Ala
248 225 230 235 240
250 Ser Leu Gly Ile Met Val Phe Val Cys Arg Asp Arg Ser Ser Ser Pro
251 245 250 255
257 <210> SEQ ID NO: 7
258 <211> LENGTH: 797
259 <212> TYPE: DNA
260 <213> ORGANISM: Viscum album coloratum
261 <221> NAME/KEY: misc_feature
262 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
264 <400> SEQUENCE: 7
265 acgatgtaac ctgcactact tccgaaccta cggtacggtt tgtgggtcga aatggcctgt 60
266 gtctcgacgt cccagagggc gattaccacg atggaagtcg gatacagttg tggccctgca 120
267 agtccaactc cgatcagaat cagctgtgga cgatcagaag ggatggaacc attcgatcta 180

```

no Xaa's
in this
sequence

(delete
circled
item)

delete circled
item
(no n's
in this
sequence)

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/627,165

DATE: 04/16/2001
TIME: 11:22:51

Input Set : A:\CRFascidoc.txt.txt
Output Set : N:\CRF3\04162001\I627165.raw

```

268 atggaaggtg cttgacgacc tatgggtata ctgcgggcag ctatataatg atctacgact 240
269 gtaatagagg ggggtgggac cttactactt ggcagataag gggcaatgga atcatcctta 300
270 atccaagatc catgatggtg atcggaacac catccgggag ccgcggaacc cgtggcacta 360
271 cttttactct gcaaacactg ggttactcat taggacaggg ctggcttgcc agcaatgata 420
272 ccgctcctcg cgaggtaacc atatatggtt tccgcgatca ttgcatggaa actagtggag 480
273 ggaaagtgtg ggttgggact tgtgtgagtg gcaagcagaa ccaaagatgg gctttgtacg 540
274 gggatgggtc cattcgcccg aaaccttacc aagaccaatg cctcacctct caggagact 600
275 ccgttagatc cgtaataaat ttatttagct gcaccgctgg atcgccaagg caacgatggg 660
276 tatttaccaa taaaggggcc attttgaatt taaagaatag gttggccatg gatgtggcgg 720
277 aatcaaatcc aagcctccgc cgaataatca tcttttcagt cactggaaat ccaaatcaaa 780
278 tgtggcttcc cgtgcca
281 <210> SEQ ID NO: 8
282 <211> LENGTH: 266
283 <212> TYPE: PRT
284 <213> ORGANISM: Viscum album coloratum
285 <221> NAME/KEY: misc_feature
286 <223> OTHER INFORMATION: any set containing "n"=X, X=any amino acid
288 <400> SEQUENCE: 8
289 Asp Asp Val Thr Cys Thr Thr Ser Glu Pro Thr Val Arg Phe Val Gly
290 1 5 10 15
292 Arg Asn Gly Leu Cys Leu Asp Val Pro Glu Gly Asp Tyr His Asp Gly
293 20 25 30
295 Ser Arg Ile Gln Leu Trp Pro Cys Lys Ser Asn Ser Asp Gln Asn Gln
296 35 40 45
298 Leu Trp Thr Ile Arg Arg Asp Gly Thr Ile Arg Ser Asn Gly Arg Cys
299 50 55 60
301 Leu Thr Thr Tyr Gly Tyr Thr Ala Gly Ser Tyr Ile Met Ile Tyr Asp
302 65 70 75 80
304 Cys Asn Arg Gly Gly Trp Asp Leu Thr Thr Trp Gln Ile Arg Gly Asn
305 85 90 95
307 Gly Ile Ile Leu Asn Pro Arg Ser Met Met Val Ile Gly Thr Pro Ser
308 100 105 110
310 Gly Ser Arg Gly Thr Arg Gly Thr Phe Thr Leu Gln Thr Leu Gly
311 115 120 125
313 Tyr Ser Leu Gly Gln Gly Trp Leu Ala Ser Asn Asp Thr Ala Pro Arg
314 130 135 140
316 Glu Val Thr Ile Tyr Gly Phe Arg Asp His Cys Met Glu Thr Ser Gly
317 145 150 155 160
319 Gly Lys Val Trp Val Gly Thr Cys Val Ser Gly Lys Gln Asn Gln Arg
320 165 170 175
322 Trp Ala Leu Tyr Gly Asp Gly Ser Ile Arg Pro Lys Pro Tyr Gln Asp
323 180 185 190
325 Gln Cys Leu Thr Ser Gln Gly Asp Ser Val Arg Ser Val Ile Asn Leu
326 195 200 205
328 Phe Ser Cys Thr Ala Gly Ser Pro Arg Gln Arg Trp Val Phe Thr Asn
329 210 215 220
331 Lys Gly Ala Ile Leu Asn Leu Lys Asn Arg Leu Ala Met Asp Val Ala
332 225 230 235 240
334 Glu Ser Asn Pro Ser Leu Arg Arg Ile Ile Ile Phe Ser Val Thr Gly

```

No Xaa's in this sequence - delete circled item

FYI!

When describing "n,"

n must represent a single nucleotide.

Please correct subsequent sequences showing similar errors

Please Note:

Use of n and/ r Xaa hav been detected in the Sequence Listing. Pleas review the Sequenc Listing to ensure that a c rresp nding xplanation is presented in the <220> t <223> fields f each sequence which presents at least ne n or Xaa.

4/16/01

VERIFICATION SUMMARY

DATE: 04/16/2001

PATENT APPLICATION: US/09/627,165

TIME: 11:22:52

Input Set : A:\CRFascidoc.txt.txt

Output Set: N:\CRF3\04162001\I627165.raw

L:21 M:283 W: Missing Blank Line separator, <400> field identifier
L:167 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:4
L:167 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:4
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:487 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
L:487 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:517 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13
L:517 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:523 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13
L:523 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:527 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:13
L:527 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:527 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:539 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:539 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:539 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:548 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:548 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:557 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:557 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:557 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14